



| The European Synchrotron



Outline

- **Introduction to BLISS**
- **LIMA Control & DAQ**
- **Scanning and Synchronization**

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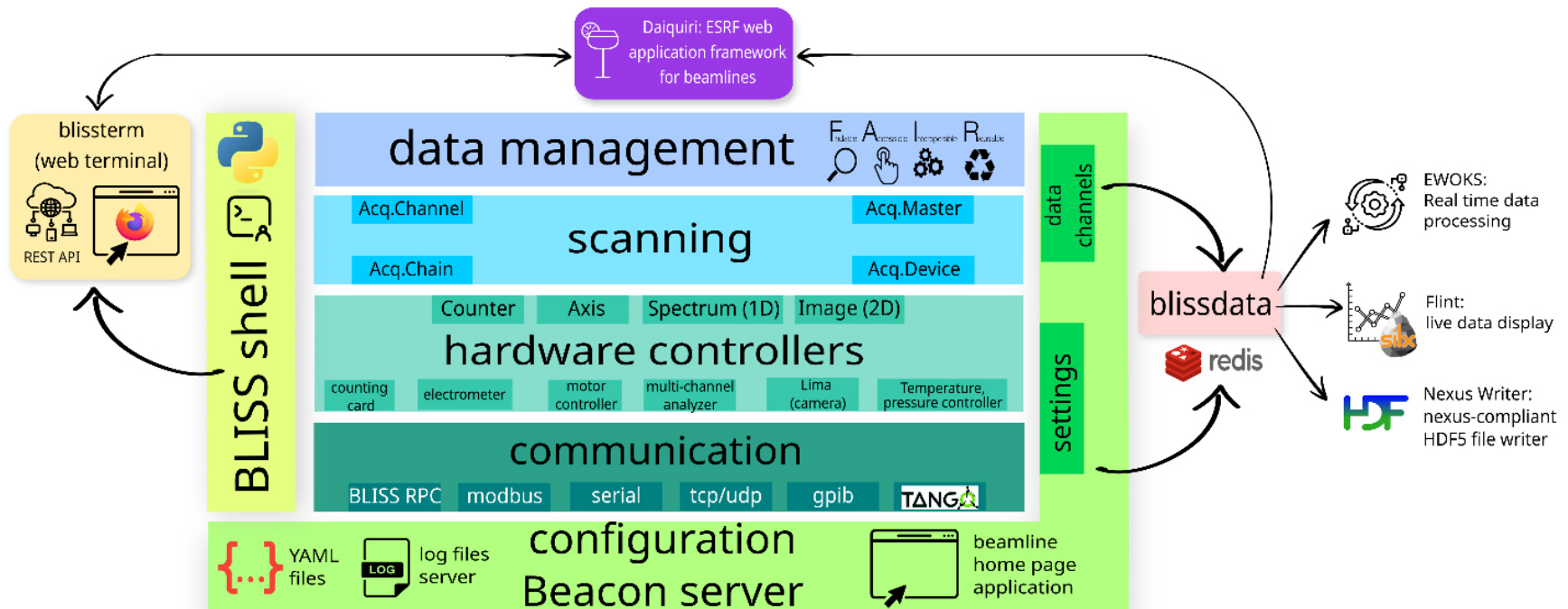
Command line driven data acquisition sequencer written in Python

Main concepts

- **Hardware abstraction layer** for all instrumentation used during a data acquisition sequence
Motors, counters, monochromators, spectrometers, diffractometers, 2D detectors, etc.
- **A generic scan engine** for step and continuous scans
The use of trajectories and HKL space is possible with all scans
- **Decoupling of data acquisition** from data saving and analysis
All data buffered in memory. Allows higher acquisition speed without blocking
- **Coherent HDF5 storage** of all acquired data at high speed and for large data volumes
All data of a proposal, its samples and the produced datasets is saved as a coherent HDF5 data tree
- **Live data display** of all acquired data
Immediate visibility of acquisition results for the user
- **Easy configuration** of hardware and experimental environment
Switch between predefined acquisitions set-ups on the fly
- **PyTango** to interface any device from the Tango world
<https://www.tango-controls.org>

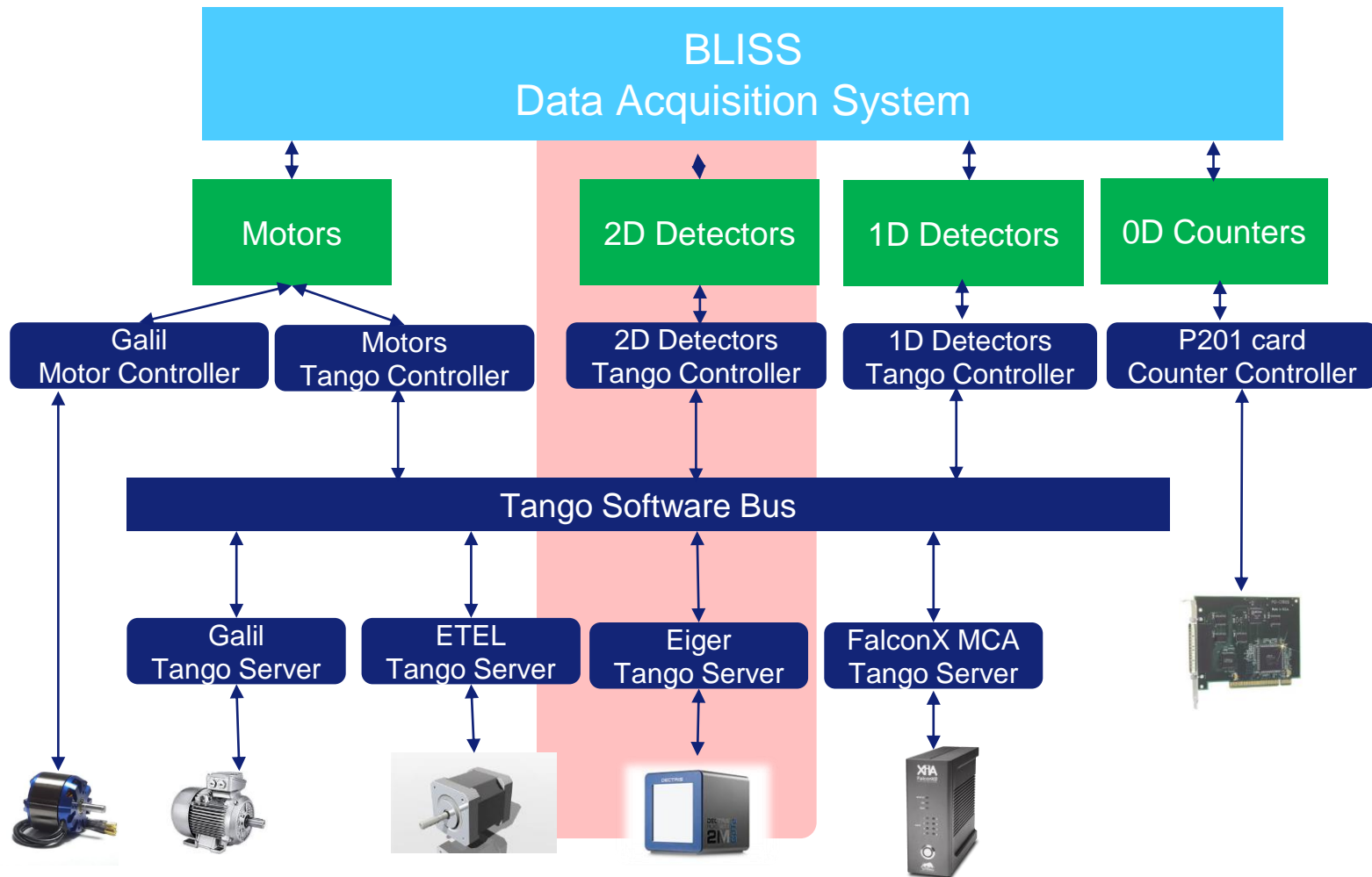
<https://bliss.gitlab-pages.esrf.fr/bliss/master>

Schematic view of the BLISS software layers



Direct hardware access or underlying control system?

All 2D detectors integrated with LIMA and wrapped as Tango Server



BLISS provides easy configuration of LIMA subsystems

- Image transformation
- Acquisition mode / accumulation
- Pixel mask
- Background subtraction
- Flat-field correction

```
TEST_EIGER [6]: deg_psi_eiger500k_1
Out [6]: deg_psi_eiger500k_1 - PSI/Eiger-500k (SlsDetector) - Lima SlsDetectorEiger

Image:
size (w,h): 1030, 514 full size (1030, 514)
depth: 1
bpp: Bpp8
binning: [1, 1]
flip: [False, False]
rotation: 0
roi: None

Acquisition:
status: |'Ready'|
status_fault_error: |'No error'|
tag: 4294967295
mode: 'SINGLE'
nb_frames: 45000
expo_time: 1e-05
trigger_mode: 'INTERNAL_TRIGGER'

Mask
----
use mask: False
mask image path: ** UNSET **
```

- Saving parameters (including expert settings):

```
      Saving
-----
File Format:  HDF5BS
L-> Suffix:  .h5
Current Mode: ONE_FILE_PER_N_FRAMES
Available Modes:
- ONE_FILE_PER_FRAME
- ONE_FILE_PER_SCAN
- ONE_FILE_PER_N_FRAMES
- SPECIFY_MAX_FILE_SIZE

for ONE_FILE_PER_N_FRAMES mode
-----
frames_per_file: 10000

for SPECIFY_MAX_FILE_SIZE mode
-----
max file size (MB): 500
L-> frames per file: 990

Expert Settings
-----
config max_writing_tasks: 1
current max_writing_tasks: 1
lima managed_mode:      SOFTWARE
```

Saving location is defined by the high-level ESRF data policy

- **ROI counter & ROI-2-Spectrum selection in Flint**

The screenshot displays the Flint software interface. The main window shows a live scan of an image titled "limatake 0.0000 45000 (#2) (finished)" with dimensions 1030 x 514. The image contains the word "EIGER" in a stylized font. Three regions of interest (ROIs) are defined: "roi1" (a rectangle), "roi2" (an arc), and "roi3" (a vertical reduction). The ROI selection panel on the right lists these ROIs with their respective kinds and checkboxes. The "roi3" entry is highlighted in blue. The "Apply these ROIs" button is visible at the bottom right of the ROI selection panel.

Flint (PID=2277) - attached to 'test_eiger'@l-cb184-1

File Windows Settings Layout Workspace Help

Live scan [base]

limatake 0.0000 45000 (#2)

sat mask stat flat expo dark time

Device deg_psi_eiger500k_1
Channel deg_psi_eiger500k_1:image
Image size 1030 x 514
Image dtype uint8

limatake 0.0000 45000 (#2) (finished)
image: 1030 x 514, id = 44999 [memory]

ROI selection

Label	Kind	
<input checked="" type="checkbox"/> roi1	Rectangle	<input type="checkbox"/>
<input checked="" type="checkbox"/> roi2	Arc	<input type="checkbox"/>
<input checked="" type="checkbox"/> roi3	Vertical reduction	<input type="checkbox"/>

Apply these ROIs

- Online data reduction:

```
ROI Counters: default
Name                Parameters                State
-----
roi1                <42,84> <537 x 178>        Enabled
roi2                <735.5, 295.1> <127.2, 183.2> <-1.7, 126.9> Enabled

Roi Profile Counters: default
Name                <x, y> <w, h> <mode>        State
-----
roi3                <17,384> <585 x 98> <horizontal> Enabled

ROI Collection: default
*** no ROI defined ***

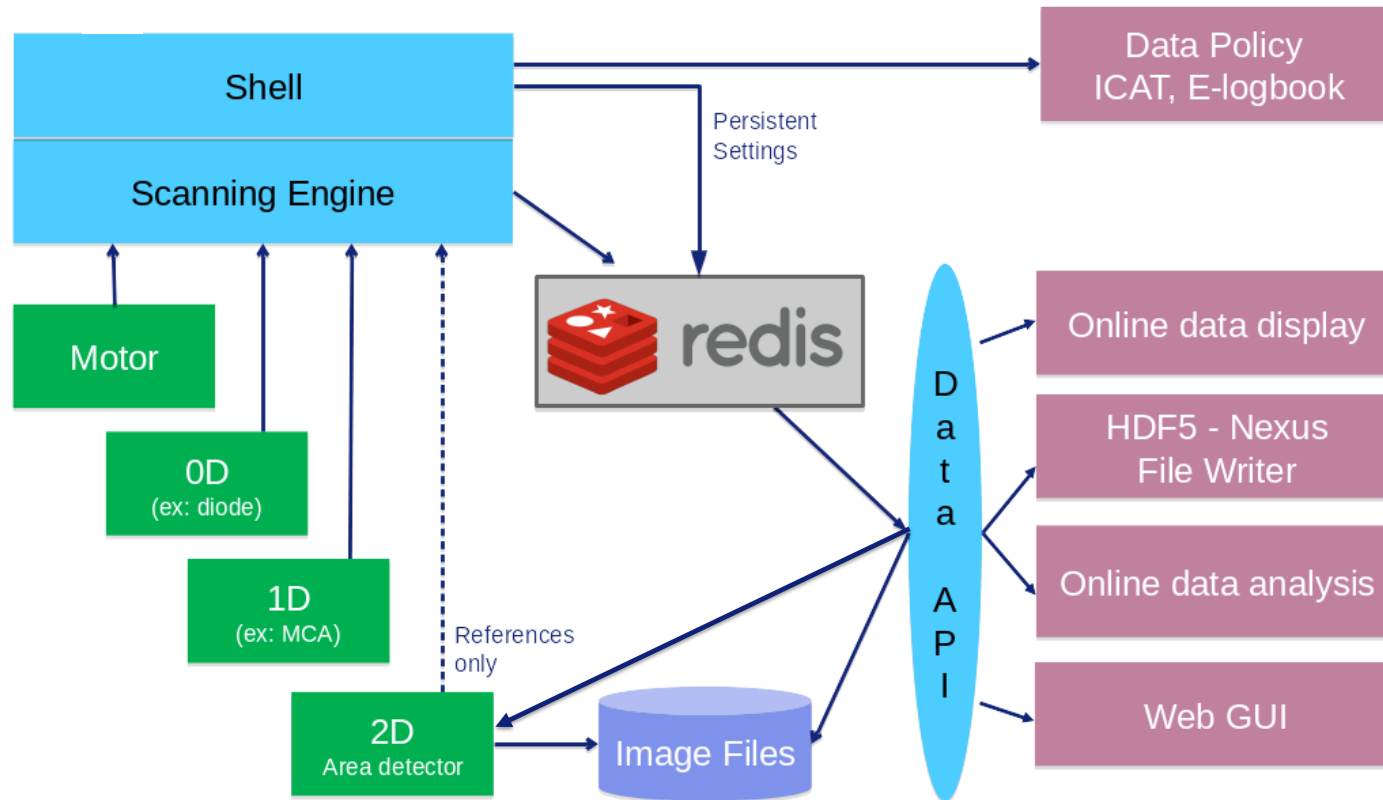
BPM Counters:
acq_time, intensity, x, y, fwhm_x, fwhm_y
```

- Detector specific parameters:

```
TEST_EIGER [7]: deg_psi_eiger500k_1.camera
Out [7]: apply_corrections: False
signed_image_mode: False
threshold_energy: -1
all_trim_bits: array([-1, -1], dtype=int32)
high_voltage: 150
clock_div: 'FULL_SPEED'
fixed_clock_div: False
parallel_mode: 'PARALLEL'
skip_frame_freq: 0
tx_frame_delay: 0
```

- And more!

Decoupling of data acquisition from data saving and analysis



- LIMA 0D & 1D data is directly injected into Redis
- Only References for 2D data
 - Reading “warm frames” from LIMA server memory
 - Reading “cold frames” from file when LIMA memory is overwritten

A high-level interface to the data in the BLISS ecosystem

- Data is injected from the source through streams
- Client register to streams and are updated as soon as new data arrives

```
scan = load a scan from blissdata
lima_stream = scan.streams["lima:image"] # Lima_stream

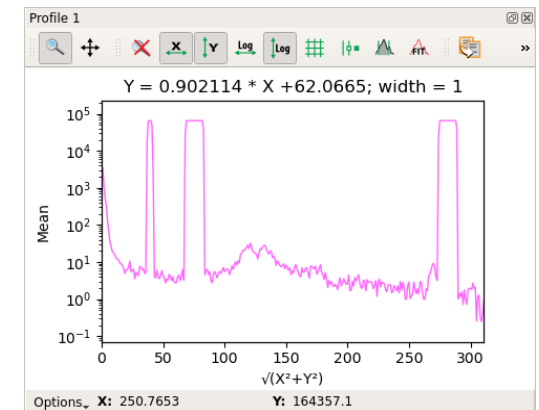
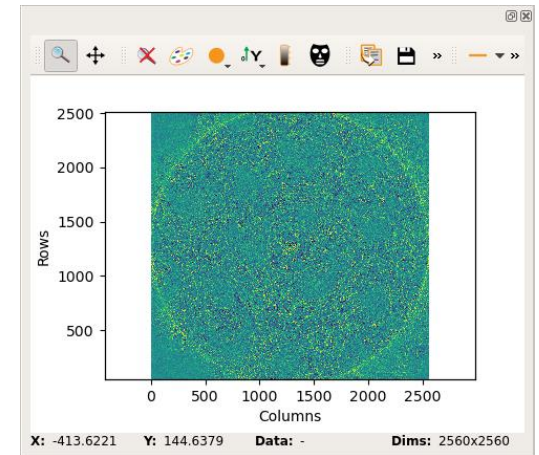
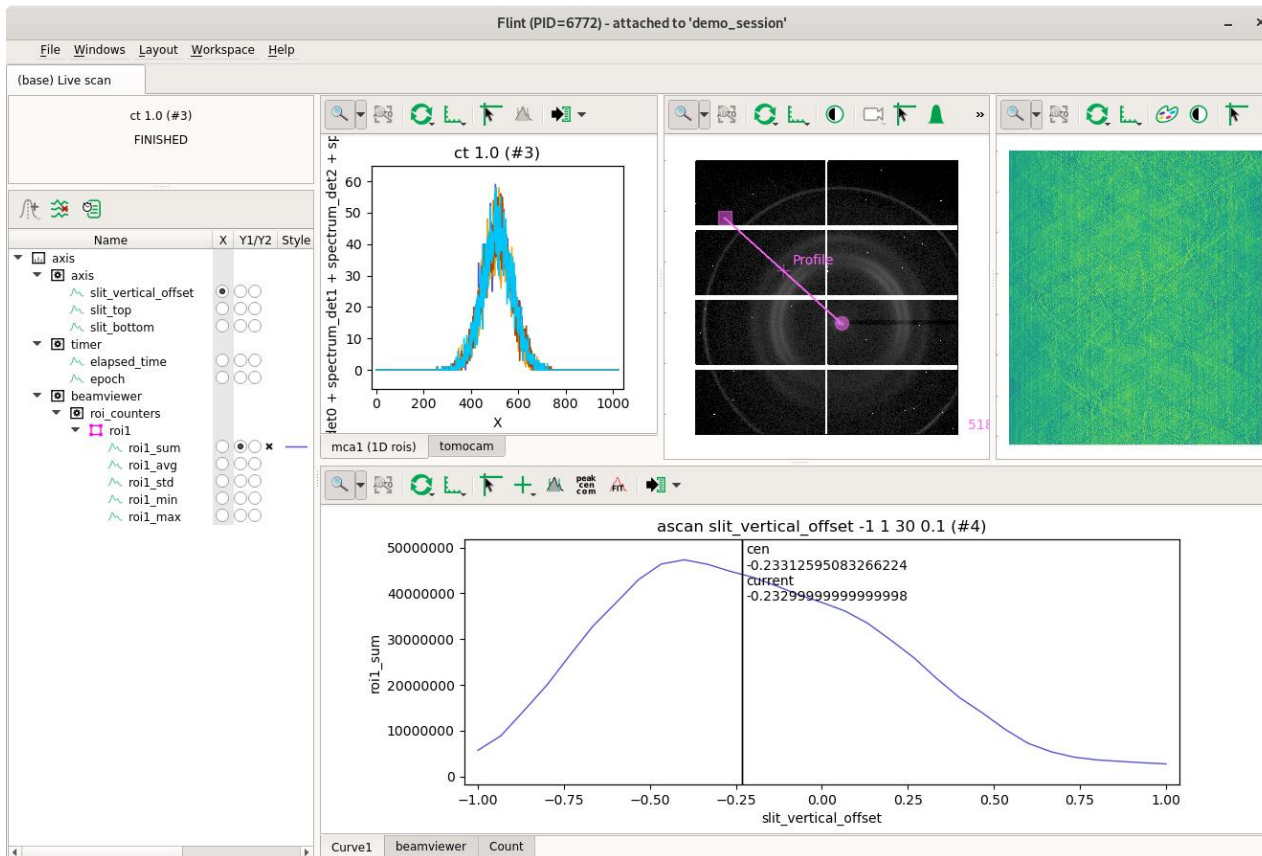
# access by index directly
try:
    lima_stream[45]
except IndexErrorNotYetThere:
    return # image 45 is not yet acquired

# access available data sequentially
cursor = lima_stream.cursor()
while True:
    try:
        view = cursor.read()
    except EndOfStream:
        return # nothing more to read
    images = view.get_data() # download all the new images
```

Flint - Live Data Display

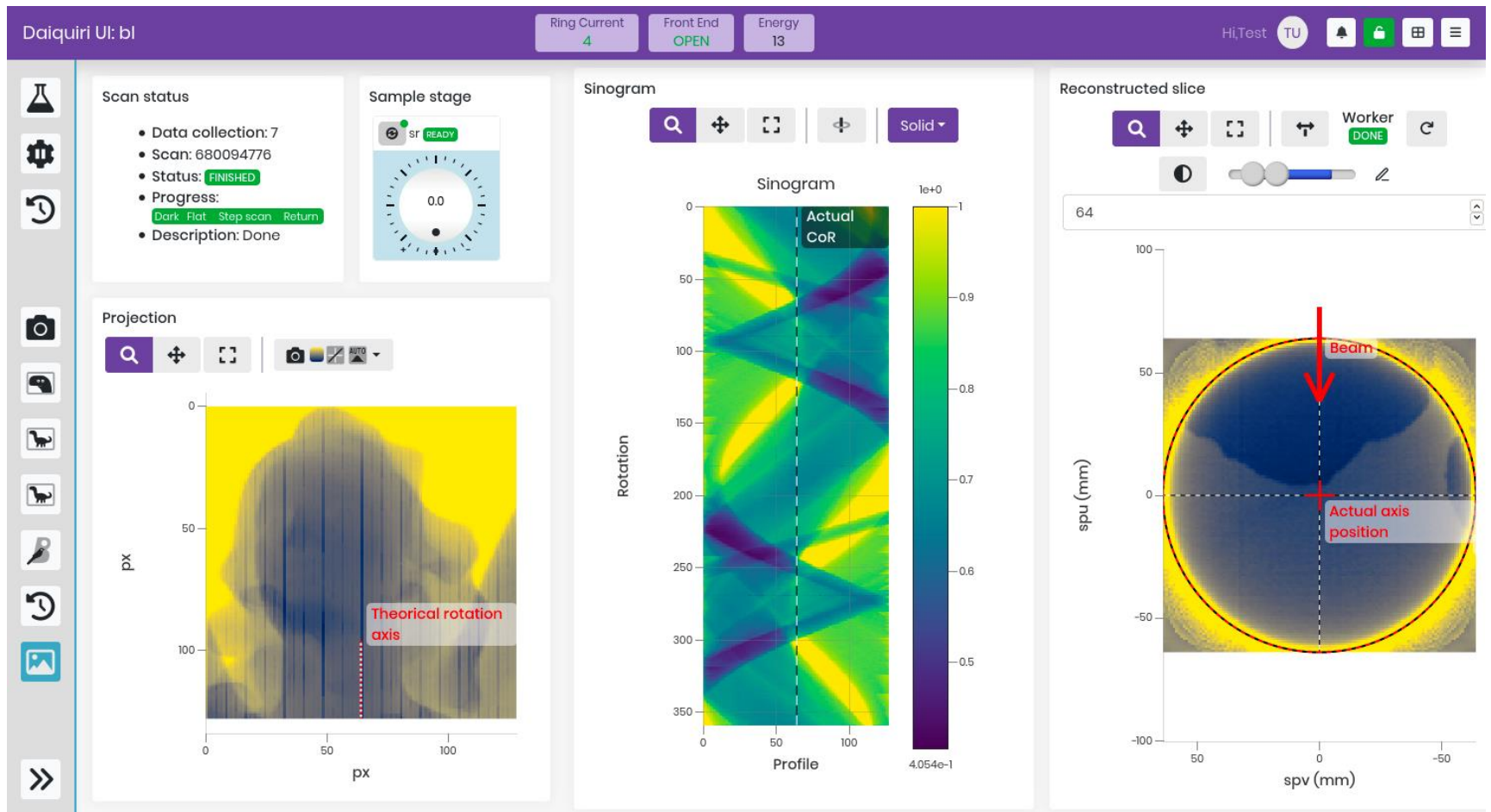
Live scan data

Calculated data can be pushed to Flint

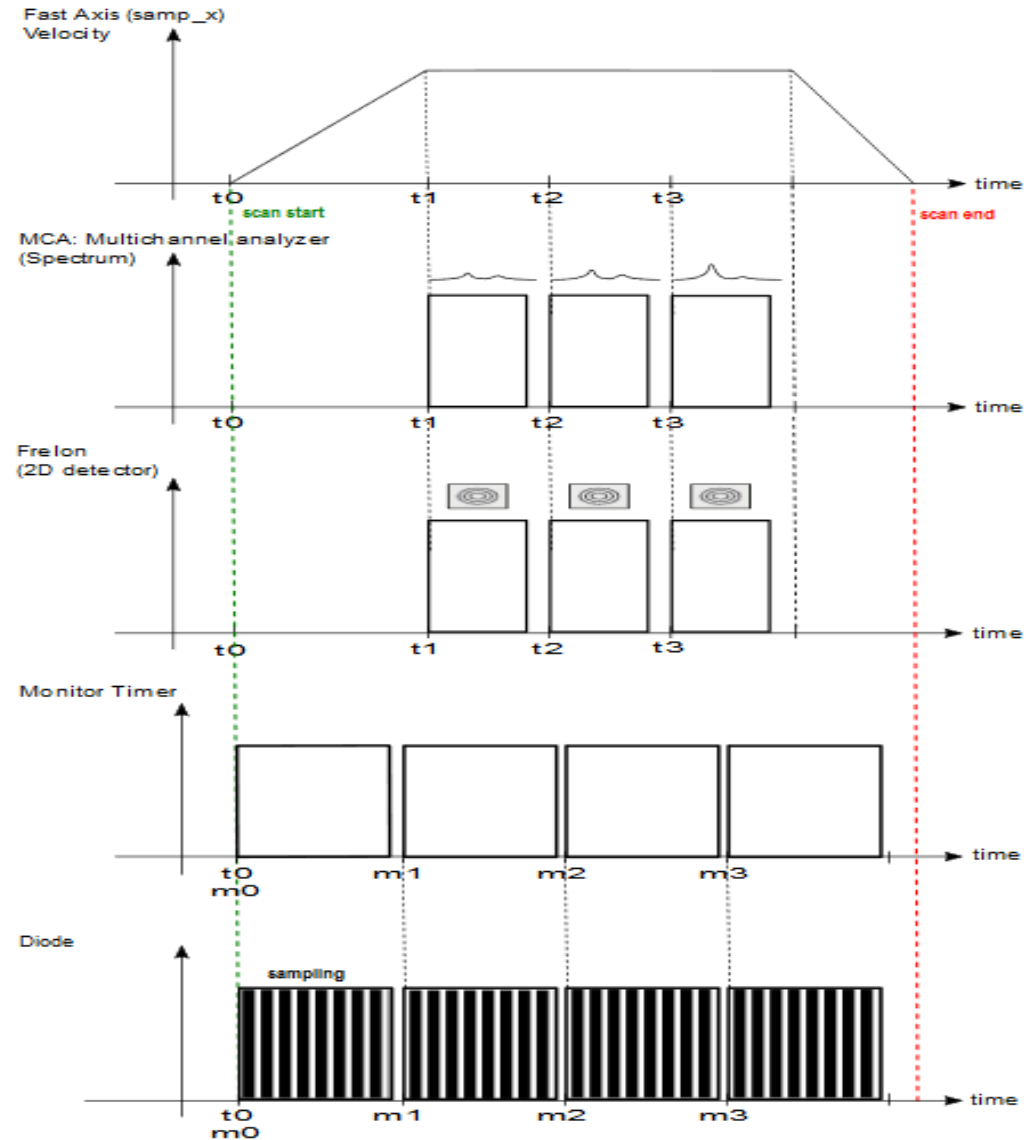
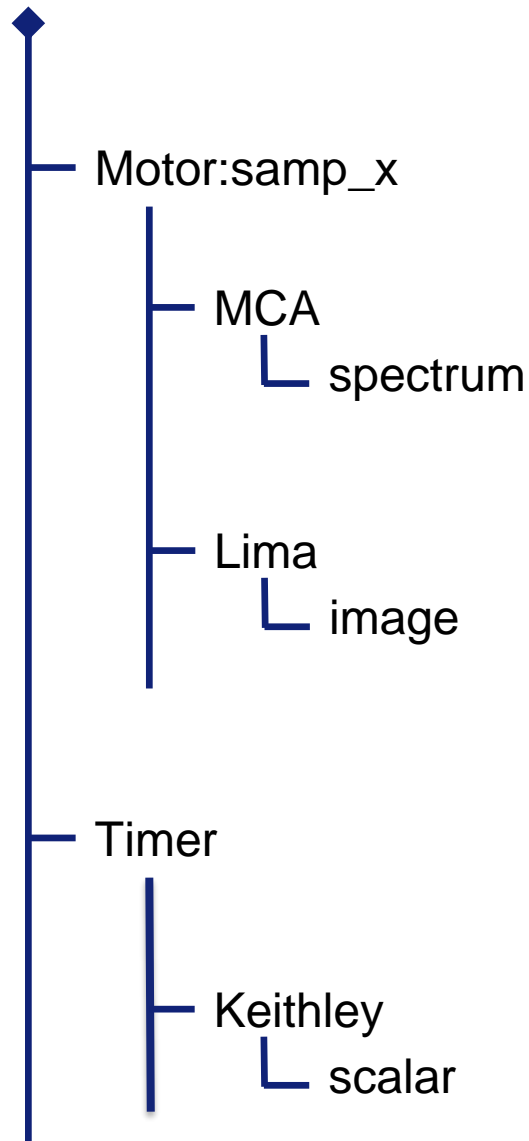


Daiquiri for Tomography

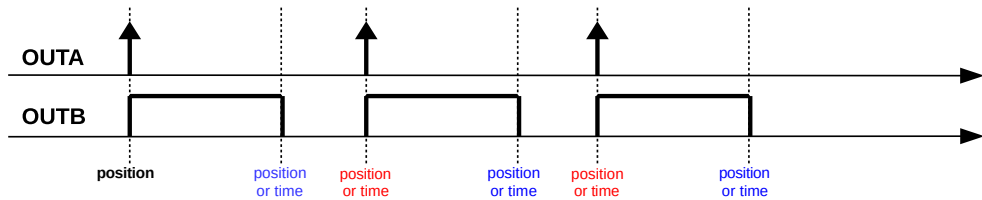
Tomography sinograms are built from Roi-2-Spectrum 1D data



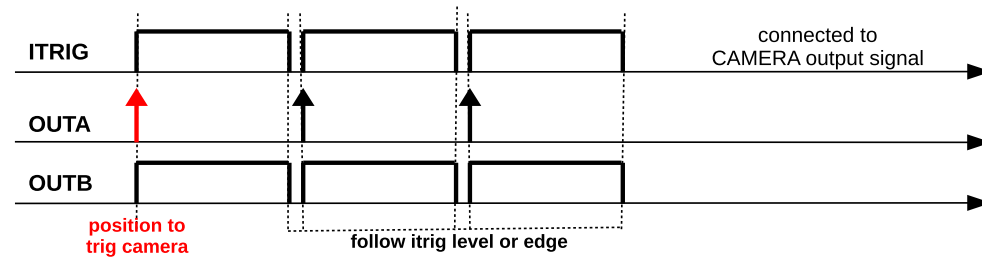
BUILDING A CONTINUOUS SCAN



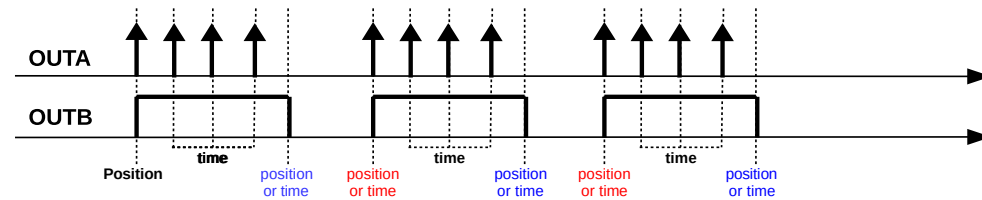
TIME or POSITION mode



CAMERA mode



LIMA accumulation supported in all modes



- **The LIMA interface is powerful but complex**
- **A high-level client library is missing for LIMA1**
- **BLISS fills this gap to make its features easy to use**
- **With LIMA2 a more complete high-level client library is under development**



Acknowledgements to

- The members of the software group for the development of all the different software tools
- Any Questions?